

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/579,648
Source: IFWP
Date Processed by STIC: 5/30/06

ENTERED

CRF Errors Edited by the STIC Systems Branch

Serial Number: 10/579,648

CRF Edit Date: 5/30/06
Edited by: h

___ Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

___ Corrected the SEQ ID NO. Sequence numbers edited were:

___ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

/ Deleted: / invalid beginning/end-of-file text ; ___ page numbers

___ Inserted mandatory headings/numeric identifiers, specifically:

___ Moved responses to same line as heading/numeric identifier, specifically:

___ Other:



IFWP

RAW SEQUENCE LISTING

DATE: 05/30/2006

PATENT APPLICATION: US/10/579,648

TIME: 15:45:56

Input Set : N:\AMC\PTO.AMC.txt

Output Set: N:\CRF4\05302006\J579648.raw

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4 <110> APPLICANT: BASF AKTIENGESELLSCHAFT et al.
6 <120> TITLE OF INVENTION: METHODS FOR THE PREPARATION OF A FINE
7   CHEMICAL BY FERMENTATION
10 <130> FILE REFERENCE: BGI-160PC2
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/579,648
C--> 12 <141> CURRENT FILING DATE: 2006-05-18
12 <150> PRIOR APPLICATION NUMBER: PCT/IB2003/006435
13 <151> PRIOR FILING DATE: 2003-12-18
15 <160> NUMBER OF SEQ ID NOS: 15
17 <170> SOFTWARE: FastSEQ for Windows Version 4.0
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 1660
21 <212> TYPE: DNA
22 <213> ORGANISM: Corynebacterium glutamicum
24 <220> FEATURE:
25 <221> NAME/KEY: CDS
26 <222> LOCATION: (301)...(1563)
28 <400> SEQUENCE: 1
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31 gcggtaccgc acccattatc gcagcatggc tggtcgaaat ctccggcgga caatggtggc 180
32 caatcgccgt ctacgtcgct gcatgttgcc ttctctctgt gatcgctcgc ttcttcaccc 240
33 aacgcgtcgc gcaccaagag aactaaaatc taagtaaaac ccctccgaaa ggaaccaccc 300
34 atg gtg aaa cgt caa ctg ccc aac ccc gca gaa cta ctc gaa ctc atg 348
35 Met Val Lys Arg Gln Leu Pro Asn Pro Ala Glu Leu Leu Glu Leu Met
36 1 5 10 15
38 aag ttc aaa aag cca gag ctc aac ggc aag aaa cga cgc cta gac tcc 396
39 Lys Phe Lys Lys Pro Glu Leu Asn Gly Lys Lys Arg Arg Leu Asp Ser
40 20 25 30
42 gcg ctc acc atc tac gac ctg cgt aaa att gct aaa cga cgc acc cca 444
43 Ala Leu Thr Ile Tyr Asp Leu Arg Lys Ile Ala Lys Arg Arg Thr Pro
44 35 40 45
46 gct gcc gcg ttc gac tac acc gac ggc gca gcc gag gcc gaa ctc tca 492
47 Ala Ala Ala Phe Asp Tyr Thr Asp Gly Ala Ala Glu Ala Glu Leu Ser
48 50 55 60
50 atc aca cgc gca cgt gaa gca ttc gaa aac atc gaa ttc cac cca gac 540
51 Ile Thr Arg Ala Arg Glu Ala Phe Glu Asn Ile Glu Phe His Pro Asp
52 65 70 75 80
54 atc ctc aag cct gca gaa cac gta gac acc acc acc caa atc ctg ggc 588
55 Ile Leu Lys Pro Ala Glu His Val Asp Thr Thr Thr Gln Ile Leu Gly
56 85 90 95
58 gga acc tcc tcc atg cca ttc ggc atc gca cca acc ggc ttc acc cgc 636
59 Gly Thr Ser Ser Met Pro Phe Gly Ile Ala Pro Thr Gly Phe Thr Arg

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RAW SEQUENCE LISTING

DATE: 05/30/2006

PATENT APPLICATION: US/10/579,648

TIME: 15:45:57

Input Set : N:\AMC\PTO.AMC.txt

Output Set : N:\CRF4\05302006\J579648.raw

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63	Leu	Met	Gln	Thr	Glu	Gly	Glu	Ile
64			115				120	
66	gca	gga	att	cct	ttc	acc	ctg	tcc
67	Ala	Gly	Ile	Pro	Phe	Thr	Leu	Ser
68		130					135	
70	gac	gtc	aag	gcc	acc	aac	ccc	aac
71	Asp	Val	Lys	Ala	Thr	Asn	Pro	Asn
72	145						150	
74	gtc	atg	cgc	gac	cgc	gaa	atc	tcc
75	Val	Met	Arg	Asp	Arg	Glu	Ile	Ser
76							165	
78	aaa	gca	gga	ttc	gac	acc	ctg	atg
79	Lys	Ala	Gly	Phe	Asp	Thr	Leu	Met
80			180				185	
82	ggc	tac	cgc	atc	cgc	gat	tcc	cgc
83	Gly	Tyr	Arg	Ile	Arg	Asp	Ser	Arg
84			195				200	
86	ctg	acc	cca	tcc	acc	gtg	ctc	aat
87	Leu	Thr	Pro	Ser	Thr	Val	Leu	Asn
88		210					215	
90	atc	gac	ttc	ctg	acc	acc	cca	acc
91	Ile	Asp	Phe	Leu	Thr	Thr	Pro	Thr
92	225						230	
94	acc	ggc	gga	acc	gtg	ggc	gac	ctc
95	Thr	Gly	Gly	Thr	Val	Gly	Asp	Leu
96							245	
98	att	tct	tac	gaa	gac	ctc	aag	gtc
99	Ile	Ser	Tyr	Glu	Asp	Leu	Lys	Val
100			260				265	
102	ctc	gta	gtc	aag	ggt	gtc	cag	aac
103	Leu	Val	Val	Lys	Gly	Val	Gln	Asn
104			275				280	
106	gac	caa	ggc	gtc	gac	ggc	ctc	atc
107	Asp	Gln	Gly	Val	Asp	Gly	Leu	Ile
108		290					295	
110	ctc	gac	cgc	gca	cca	gtc	cca	ttc
111	Leu	Asp	Arg	Ala	Pro	Val	Pro	Phe
112	305						310	
114	gaa	gtc	gga	tct	gaa	cca	acc	atc
115	Glu	Val	Gly	Ser	Glu	Pro	Thr	Ile
116							325	
118	ggc	gcc	gac	atc	gtc	gca	gcc	gta
119	Gly	Ala	Asp	Ile	Val	Ala	Ala	Val
120							340	
122	atc	ggt	cgt	gcc	tac	ctc	tac	gga
123	Ile	Gly	Arg	Ala	Tyr	Leu	Tyr	Gly
124			355				360	

RAW SEQUENCE LISTING

DATE: 05/30/2006

PATENT APPLICATION: US/10/579,648

TIME: 15:45:57

Input Set : N:\AMC\PTO.AMC.txt

Output Set: N:\CRF4\05302006\J579648.raw

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127 Val Asp Arg Thr Ile Ala Ile Leu Arg Ser Glu Ile Thr Arg Thr Met
128 370 375 380
130 gct ctc ctc ggt gtt tcc tcc ctc gaa gaa ctc gag cca cgc cac gtc 1500
131 Ala Leu Leu Gly Val Ser Ser Leu Glu Glu Leu Glu Pro Arg His Val
132 385 390 395 400
134 acc cag ctg gcc aag atg gtt cca gtt tct gac gca act cgt tct gca 1548
135 Thr Gln Leu Ala Lys Met Val Pro Val Ser Asp Ala Thr Arg Ser Ala
136 405 410 415
138 gcg gcg gag att taa aagtttctct ccttagctat taaaagggtgc ccatccgttt 1603
139 Ala Ala Glu Ile *
140 420
142 ggatgggcac cttctcgttt cttgcaatcg gcatattcag tcaaaaaatg ttgaaat 1660
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145 <211> LENGTH: 420
146 <212> TYPE: PRT
147 <213> ORGANISM: Corynebacterium glutamicum
149 <400> SEQUENCE: 2
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151 1 5 10 15
152 Lys Phe Lys Lys Pro Glu Leu Asn Gly Lys Lys Arg Arg Leu Asp Ser
153 20 25 30
154 Ala Leu Thr Ile Tyr Asp Leu Arg Lys Ile Ala Lys Arg Arg Thr Pro
155 35 40 45
156 Ala Ala Ala Phe Asp Tyr Thr Asp Gly Ala Ala Glu Ala Glu Leu Ser
157 50 55 60
158 Ile Thr Arg Ala Arg Glu Ala Phe Glu Asn Ile Glu Phe His Pro Asp
159 65 70 75 80
160 Ile Leu Lys Pro Ala Glu His Val Asp Thr Thr Thr Gln Ile Leu Gly
161 85 90 95
162 Gly Thr Ser Ser Met Pro Phe Gly Ile Ala Pro Thr Gly Phe Thr Arg
163 100 105 110
164 Leu Met Gln Thr Glu Gly Glu Ile Ala Gly Ala Gly Ala Ala Gly Ala
165 115 120 125
166 Ala Gly Ile Pro Phe Thr Leu Ser Thr Leu Gly Thr Thr Ser Ile Glu
167 130 135 140
168 Asp Val Lys Ala Thr Asn Pro Asn Gly Arg Asn Trp Phe Gln Leu Tyr
169 145 150 155 160
170 Val Met Arg Asp Arg Glu Ile Ser Tyr Gly Leu Val Glu Arg Ala Ala
171 165 170 175
172 Lys Ala Gly Phe Asp Thr Leu Met Phe Thr Val Asp Thr Pro Ile Ala
173 180 185 190
174 Gly Tyr Arg Ile Arg Asp Ser Arg Asn Gly Phe Ser Ile Pro Pro Gln
175 195 200 205
176 Leu Thr Pro Ser Thr Val Leu Asn Ala Ile Pro Arg Pro Trp Trp Trp
177 210 215 220
178 Ile Asp Phe Leu Thr Thr Pro Thr Leu Glu Phe Ala Ser Leu Ser Ser
179 225 230 235 240
180 Thr Gly Gly Thr Val Gly Asp Leu Leu Asn Ser Ala Met Asp Pro Thr

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DATE: 05/30/2006

PATENT APPLICATION: US/10/579,648

TIME: 15:45:57

Input Set : N:\AMC\PTO.AMC.txt

Output Set: N:\CRF4\05302006\J579648.raw

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181          245          250          255
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183          260          265          270
184 Leu Val Val Lys Gly Val Gln Asn Val Glu Asp Ser Val Lys Leu Leu
185          275          280          285
186 Asp Gln Gly Val Asp Gly Leu Ile Leu Ser Asn His Gly Gly Arg Gln
187          290          295          300
188 Leu Asp Arg Ala Pro Val Pro Phe His Leu Leu Pro Gln Val Arg Lys
189 305          310          315          320
190 Glu Val Gly Ser Glu Pro Thr Ile Met Ile Asp Thr Gly Ile Met Asn
191          325          330          335
192 Gly Ala Asp Ile Val Ala Ala Val Ala Met Gly Ala Asp Phe Thr Leu
193          340          345          350
194 Ile Gly Arg Ala Tyr Leu Tyr Gly Leu Met Ala Gly Gly Arg Glu Gly
195          355          360          365
196 Val Asp Arg Thr Ile Ala Ile Leu Arg Ser Glu Ile Thr Arg Thr Met
197          370          375          380
198 Ala Leu Leu Gly Val Ser Ser Leu Glu Glu Leu Glu Pro Arg His Val
199 385          390          395          400
200 Thr Gln Leu Ala Lys Met Val Pro Val Ser Asp Ala Thr Arg Ser Ala
201          405          410          415
202 Ala Ala Glu Ile
203          420
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207 <211> LENGTH: 35
208 <212> TYPE: DNA
209 <213> ORGANISM: Artificial Sequence
211 <220> FEATURE:
212 <223> OTHER INFORMATION: Oligonucleotide
214 <400> SEQUENCE: 3
215 gagagagaga cgcgtccag tggctgagac gcatc 35
217 <210> SEQ ID NO: 4
218 <211> LENGTH: 34
219 <212> TYPE: DNA
220 <213> ORGANISM: Artificial Sequence
222 <220> FEATURE:
223 <223> OTHER INFORMATION: Oligonucleotide
225 <400> SEQUENCE: 4
226 ctctctctgt cgacgaattc aatcttacgg cctg 34
228 <210> SEQ ID NO: 5
229 <211> LENGTH: 4323
230 <212> TYPE: DNA
231 <213> ORGANISM: Corynebacterium glutamicum
233 <400> SEQUENCE: 5
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235 tatcgtcgac atcgatgctc ttctgctgta attaacaatt gggatcctct agaccggga 120
236 tttaaatcgc tagcgggctg cttaaaggaag cggaacacgt agaaagccag tccgcagaaa 180
237 cggtgctgac cccggatgaa tgtcagctac tgggctatct ggacaaggga aaacgcaagc 240
238 gcaaagagaa agcaggtagc ttgcagtggg cttacatggc gatagctaga ctgggcgggt 300

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RAW SEQUENCE LISTING

DATE: 05/30/2006

PATENT APPLICATION: US/10/579,648

TIME: 15:45:57

Input Set : N:\AMC\PTO.AMC.txt

Output Set: N:\CRF4\05302006\J579648.raw

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239 ttatggacag caagcgaacc ggaattgcca gctggggcgc cctctggtaa gggtgggaag 360
240 cctgcaaag taaactggat ggctttcttg ccgccaagga tctgatggcg caggggatca 420
241 agatctgac aagagacagg atgaggatcg ttctgcatga ttgaacaaga tggattgcac 480
242 gcaggttctc cggccgcttg ggtggagagg ctattcggct atgactgggc acaacagaca 540
243 atcggctgct ctgatgccgc cgtgttcggg ctgtcagcgc aggggcgccc ggttcttttt 600
244 gtcaagaccg acctgtccgg tgccctgaat gaactgcagg acgaggcagc gcggctatcg 660
245 tggctggcca cgacgggcgt tccttgcgca gctgtgctcg acgttgtcac tgaagcggga 720
246 agggactggc tgctattggg cgaagtgcgg gggcaggatc tcctgtcatc tcacctgtct 780
247 cctgccgaga aagtatccat catggctgat gcaatgcggc ggctgcatac gcttgatccg 840
248 gctacctgcc cattcgacca ccaagcgaaa catcgcatcg agcagcacg tactcggatg 900
249 gaagccggtc ttgtcgatca ggatgatctg gacgaagagc atcaggggct cgcgccagcc 960
250 gaactgttcg ccaggctcaa ggcgcgcgat cccgacggcg aggatctcgt cgtgacctat 1020
251 ggcgatgcct gcttgccgaa tatcatggtg gaaaatggcc gcttttcttg attcatcgac 1080
252 tgtggccggc tgggtgtggc ggaccgctat caggacatag cgttggctac ccgtgatatt 1140
253 gctgaagagc ttggcggcga atgggctgac cgcttcctcg tgctttacgg tatcgccgct 1200
254 cccgattcgc agcgcacgc cttctatcgc cttcttgacg agttcttctg agcgggactc 1260
255 tggggttcga aatgaccgac caagcgacgc ccaacctgcc atcacgagat ttcgattcca 1320
256 ccgccgcctt ctatgaaagg ttgggcttcg gaatcgtttt ccgggacgcc ggctggatga 1380
257 tcctccagcg cggggatctc atgctggagt tcttcgccc cgtagcggc gcgccggccg 1440
258 gccgggtgtg aaataccgca cagatgcgta aggagaaaat accgcacag gcgctcttcc 1500
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265 gcgctttctc atagctcacg ctgtaggatc ctgagttcgg tgtaggctgt tcgctccaag 1920
266 ctgggctgtg tgcacgaacc ccccgctcag cccgaccgct gcgccttatc cggtaactat 1980
267 cgtcttgagt ccaaccgggt aagacacgac ttatcgccac tggcagcagc cactggtaac 2040
268 aggattagca gagcgaggta tgtaggcggg gctacagagt tcttgaagtg gtggcctaac 2100
269 tacggctaca ctagaaggac agtatttggt atctgcgctc tgctgaagcc agttaccttc 2160
270 ggaaaaagag ttggtagctc ttgatccggc aaacaaacca ccgctggtag cgggtggtttt 2220
271 tttgtttgca agcagcagat tacgcgcaga aaaaaaggat ctcaagaaga tcctttgatc 2280
272 ttttctacgg ggtctgacgc tcagtggaac gaaaactcac gttaagggat tttggtcatg 2340
273 agattatcaa aaaggatctt cacctagatc cttttaaagg ccggcccgcg ccgccatcgg 2400
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283 ctttgagaa gtttttgact ttcttgacgg aagaatgatg tgcttttgcc atagtatgct 3000
284 ttgttaaata aagattcttc gccttggtag ccatcttcag ttccagtgtt tgcttcaaat 3060
285 actaagtatt tgtggccttt atcttctacg tagtgaggat ctctcagcgt atgggtgtcg 3120
286 cctgagctgt agttgccttc atcgatgaac tgctgtacat tttgatacgt tttccgtca 3180
287 ccgtcaaaag ttgatttata atcctctaca ccgttgatgt tcaaagagct gtctgatgct 3240

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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/579,648

DATE: 05/30/2006

TIME: 15:45:58

Input Set : N:\AMC\PTO.AMC.txt

Output Set: N:\CRF4\05302006\J579648.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

**Raw Sequence Listing before editing
(for reference only)**



IFWP

RAW SEQUENCE LISTING

DATE: 05/26/2006

PATENT APPLICATION: US/10/579,648

TIME: 08:25:28

Input Set : F:\SEQLIST.txt

Output Set: N:\CRF4\05262006\J579648.raw

4 <110> APPLICANT: BASF AKTIENGESELLSCHAFT et al.
 6 <120> TITLE OF INVENTION: METHODS FOR THE PREPARATION OF A FINE
 7 CHEMICAL BY FERMENTATION
 10 <130> FILE REFERENCE: BGI-160PC2
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/579,648
 C--> 12 <141> CURRENT FILING DATE: 2006-05-18
 12 <150> PRIOR APPLICATION NUMBER: PCT/IB2003/006435
 13 <151> PRIOR FILING DATE: 2003-12-18
 15 <160> NUMBER OF SEQ ID NOS: 15
 17 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

819 <210> SEQ ID NO: 15
 820 <211> LENGTH: 7561
 821 <212> TYPE: DNA
 822 <213> ORGANISM: Corynebacterium glutamicum
 824 <400> SEQUENCE: 15

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 827 gccgttttgct aactcaaccg tgcgtttttt atcgttttgc agaagttttt gactttcttg 180
 828 acggaagaat gatgtgcttt tgccatagta tgctttgtta aataaagatt cttcgccctg 240
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 839 ggacgaatca aattcagaaa cttgatattt ttcatttttt tgctgttcag ggatttgcag 900
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 848 attctctgta ttttttatag tttctgttgc atgggcataa agttgccttt ttaatcacia 1440

p.4

**Does Not Comply
Corrected Diskette Needed**

RAW SEQUENCE LISTING

DATE: 05/26/2006

PATENT APPLICATION: US/10/579,648

TIME: 08:25:29

Input Set : F:\SEQLIST.txt

Output Set: N:\CRF4\05262006\J579648.raw

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851 agaaggcgat agaaggcgat gcgctgcgaa tcgggagcgg cgataccgta aagcacgagg 1620
852 aagcggtcag cccattcgcc gccaaagctct tcagcaatat cacgggtagc caacgctatg 1680
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857 cgatgtttcg cttggtggtc gaatgggcag gtagccggat caagcgtatg cagccgccgc 1980
858 attgcatcag ccatgatgga tactttctcg gcaggagcaa ggtgagatga caggagatcc 2040
859 tgccccggca cttcgcccaa tagcagccag tcccttcccg cttcagtgac aacgtcgagc 2100
860 acagctgcgc aaggaacgcc cgtcgtggcc agccacgata gccgcgctgc ctcgctcttg 2160
861 agttcattca gggcaccgga caggtcggtc ttgacaaaaa gaaccgggcg cccctgcgct 2220
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863 aatagcctct ccaccaagc ggccggagaa cctgcgtgca atccatcttg ttcaatcatg 2340
864 cgaaacgata ctcatcctgt ctcttgatca gatcttgatc ccctgcgcca tcagatcctt 2400
865 ggccgcaaga aagccatcca gtttactttg cagggtctcc caaccttacc agaggggcgc 2460
866 ccagctggca attccggttc gcttgctgtc cataaaaccg ccagtcctag ctatcgccat 2520
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RAW SEQUENCE LISTING

DATE: 05/26/2006

PATENT APPLICATION: US/10/579,648

TIME: 08:25:29

Input Set : F:\SEQLIST.txt

Output Set: N:\CRF4\05262006\J579648.raw

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Input Set : F:\SEQLIST.txt

Output Set: N:\CRF4\05262006\J579648.raw

947 tgtctgcgta gaatcctctg tttgtcatat agcttgtaat caccgacattg tttccttytc 7380
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951 g 7561

E--> 953 Bgl-160Pc2

E--> 955 - 1 -

delete

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/579,648

DATE: 05/26/2006

TIME: 08:25:30

Input Set : F:\SEQLIST.txt

Output Set: N:\CRF4\05262006\J579648.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:953 M:254 E: No. of Bases conflict, LENGTH:Input:2 Counted:7567 SEQ:15
L:953 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:6
L:953 M:112 C: (48) String data converted to lower case,
M:254 Repeated in SeqNo=15
L:955 M:252 E: No. of Seq. differs, <211> LENGTH:Input:7561 Found:7567 SEQ:15